

10019258 .092302 ~~10/0192585~~

SEQUENCE LISTING ~~10 Rows~~ 23 SEP 2002

<110> Lexow, Preben
<120> Method of cloning and producing fragment chains with readable information content
<130> 1181-256
<140> US 10/019258
<141> 2001-12-28
<150> PCT/GB00/02512
<151> 2000-06-27
<150> NO 20003191
<151> 2000-06-20
<150> NO 20003190
<151> 2000-06-20
<150> NO 19991325
<151> 1999-06-28
<160> 105
<170> PatentIn version 3.1
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<223> N is any nucleotide.

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11

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 <223> N is any nucleotide.

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11

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 <223> BbvI overhang

<400> 3
 cgagcgccctc cagtgcagcg gag

23

<210> 4
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 <223> BbvI overhang

<400> 4
 tatcgccgcct ccagtgcagc ggag

24

<210> 5
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 <212> DNA
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<220>
 <223> BbvI overhang

<400> 5
 ctctcgccct ccagtgcagc ggag

24

<210> 6
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 <212> DNA
 <213> Artificial Sequence

<220>
 <223> BbvI overhang 6 (delC)

<400> 6
 ctctctccgc tgcactggag gcgc

24

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<223> BbvI overhang 7a		
<400> 7		
caacgcgcct ccagtgcagc ggag		24
<210> 8		
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<400> 8		
ggtaggcgcct ccagtgcagc ggag		24
<210> 9		
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aagagctccg ctgcactgga ggcgc		25
<210> 10		
<211> 25		
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<400> 10		
ctcttctccg ctgcactgga ggcgc		25
<210> 11		
<211> 35		
<212> DNA		
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<220>
 <223> Consensus binding motifs of the initiation linkers
 <220>
 <221> misc_feature
 <222> (19)..(24)
 <223> N is any nucleotide.

<400> 11
 gcagcgacca tgagtccanc tcnnngtggat gacgc

35

<210> 12
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 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Initiation linker

<220>
 <221> misc_feature
 <222> (19)..(37)
 <223> N is any nucleotide with the proviso that the DNA sequence from 3
 2 to 37 is not palindromic.

<400> 12
 gcagcgacca tgagtccanc tcnnngtggat gnnnnnn

37

<210> 13
 <211> 38
 <212> DNA
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<220>
 <223> Initiation linker

<220>
 <221> misc_feature
 <222> (19)..(38)
 <223> N is any nucleotide with the proviso that the DNA sequence from 3
 3 to 38 is not palindromic.

<400> 13
 gcagcgacca tgagtccanc tcnnngtggat gnnnnnnn

38

<210> 14
 <211> 39
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Initiation linker

 <220>
 <221> misc_feature
 <222> (19)..(39)
 <223> N is any nucleotide with the proviso that the DNA sequence from 3
 4 to 39 is not palindromic.

<400> 14
 gcagcgacca tgagtccanc tcnnngtggat gnnnnnnnnn 39

<210> 15
 <211> 40
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Initiation linker

 <220>
 <221> misc_feature
 <222> (19)..(40)
 <223> N is any nucleotide with the proviso that the DNA sequence from 3
 5 to 40 is not palindromic.

<400> 15
 gcagcgacca tgagtccanc tcnnngtggat gnnnnnnnnnnn 40

<210> 16
 <211> 41
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Initiation linker

 <220>
 <221> misc_feature
 <222> (19)..(41)
 <223> N is any nucleotide with the proviso that the DNA sequence from 3
 6 to 41 is not palindromic.

<400> 16
 gcagcgacca tgagtccanc tcnnngtggat gacgcnnnn n 41

<210> 17
 <211> 42

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<212> DNA
<213> Artificial Sequence

<220>
<223> Initiation linker

<220>
<221> misc_feature
<222> (19)..(42)
<223> N is any nucleotide with the proviso that the DNA sequence from 3
      7 to 42 is not palindromic.

<400> 17
gcagcgacca tgagtccanc tcnnngtggat gacgcnnnn nn          42

<210> 18
<211> 43
<212> DNA
<213> Artificial Sequence

<220>
<223> Initiation linker

<220>
<221> misc_feature
<222> (19)..(43)
<223> N is any nucleotide with the proviso that the DNA sequence from 3
      8 to 43 is not palindromic.

<400> 18
gcagcgacca tgagtccanc tcnnngtggat gacgcnnnn nnn          43

<210> 19
<211> 44
<212> DNA
<213> Artificial Sequence

<220>
<223> Initiation linker

<220>
<221> misc_feature
<222> (19)..(44)
<223> N is any nucleotide with the proviso that the DNA sequence from 3
      9 to 44 is not palindromic.

<400> 19
gcagcgacca tgagtccanc tcnnngtggat gacgcnnnn nnnn          44

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<210> 20
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
<223> Initiation linker

<220>
<221> misc_feature
<222> (19)..(45)
<223> N is any nucleotide with the proviso that the DNA sequence from 4
0 to 45 is not palindromic.

<400> 20
gcagcgacca tgagtccanc tcnnngtggat gacgcnnnn nnnnnn 45

<210> 21
<211> 46
<212> DNA
<213> Artificial Sequence

<220>
<223> Initiation linker

<220>
<221> misc_feature
<222> (19)..(46)
<223> N is any nucleotide with the proviso that the DNA sequence from 4
1 to 46 is not palindromic.

<400> 21
gcagcgacca tgagtccanc tcnnngtggat gacgcnnnn nnnnnn 46

<210> 22
<211> 50
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic oligonucleotide

<400> 22
taatacgact cactatacca caagtttgta caaaaaagca ggctctattc 50

<210> 23
<211> 56
<212> DNA
<213> Artificial Sequence

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<220>
<223> Synthetic oligonucleotide

<400> 23
taggaagaat agagcctgct ttttgtaca aacttgtggt atagtgagtc gtatta      56

<210> 24
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic oligonucleotide

<400> 24
ttccttatgca gtggaccact ttgtacaaga aagctgggtt gcagt      45

<210> 25
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic oligonucleotide

<400> 25
gcaactactg caacccagct ttcttgtaca aagtggtcca ctgca      45

<210> 26
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic oligonucleotide

<400> 26
agttgcttga cgccacaagt ttgtacaaaa aagcaggctt tgacg      45

<210> 27
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic oligonucleotide

<400> 27
cgacatcgtc aaagcctgct ttttgtaca aacttgtggc gtcaa      45

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<210> 28
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic oligonucleotide

<400> 28
atgtcgaagg gcggaccact ttgtacaaga aagctggta agggc          45

<210> 29
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic oligonucleotide

<400> 29
gacagggccc ttacccagct ttcttgtaca aagtggtccg ccctt          45

<210> 30
<211> 58
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic oligonucleotide

<400> 30
cctgtcatgt ggaccacttt gtacaagaaa gctgggttc tatagtgtca cctaaatc      58

<210> 31
<211> 52
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic oligonucleotide

<400> 31
gatttaggtg acactataga aacccagctt tcttgtacaa agtggtccac at          52

<210> 32
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic oligonucleotide

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<400> 32		
taatacgact cactatacca		20
<210> 33		
<211> 17		
<212> DNA		
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<223> Synthetic oligonucleotide		
<400> 33		
taatacgact cactata		17
<210> 34		
<211> 21		
<212> DNA		
<213> Artificial Sequence		
<220>		
<223> Synthetic oligonucleotide		
<400> 34		
aagatatatcac agtggattta g		21
<210> 35		
<211> 21		
<212> DNA		
<213> Artificial Sequence		
<220>		
<223> Fragment chain 2 terminal		
<400> 35		
ttctatacgatgc tcacctaataat c		21
<210> 36		
<211> 46		
<212> DNA		
<213> Artificial Sequence		
<220>		
<223> Primer		
<400> 36		
tcaacggcaa cctacatgac catccgattt aggtgacact atagaa		46
<210> 37		
<211> 47		

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<212> DNA
<213> Artificial Sequence

<220>
<223> Primer

<400> 37
gtcatgttagg ttgccgttga tccatcctaa tacgactcac tatacgat 47

<210> 38
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Fragment chain 3 terminal

<400> 38
tgctatagtg agtcgtattat 20

<210> 39
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Initiation linker 1 (s)

<400> 39
attcggtcga gatgctctca 20

<210> 40
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Initiation linker 1 (as)

<400> 40
cgactgagag catctcgacc gaat 24

<210> 41
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Initiation linker 2

<400> 41

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gcgttactga gcgtagctct g

21

<210> 42
 <211> 25
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Initiation linker 2 (as)

 <400> 42
 ctctcagagc tacgctcagt aacgc

25

<210> 43
 <211> 24
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Propagation linker (s)

 <220>
 <221> misc_feature
 <222> (20)..(24)
 <223> N is any nucleotide.

<400> 43
 tgctgcagga gcgaatctcn nnnn

24

<210> 44
 <211> 19
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Propagation linker (as)

 <400> 44
 gagattcgct cctgcagca

19

<210> 45
 <211> 24
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Labeling linker 2 (s)

 <400> 45
 ctcttgctat agtgagtcgt atta

24

<210> 46
 <211> 20
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Labeling linker 2 (as)

<400> 46
 taatacgact cactatacgca

20

<210> 47
 <211> 30
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Termination linker 1 (s)

<400> 47
 aagagctcag gtcattgacg tagctatgaa

30

<210> 48
 <211> 20
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Termination linker 1/2 (as)

<400> 48
 agctacgtca atgacacctgag

20

<210> 49
 <211> 10
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Termination linker 1 (short version)

<400> 49
 aagagatgaa

10

<210> 50
 <211> 29
 <212> DNA
 <213> Artificial Sequence

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<220>
<223> Termination linker 2 (s)

<400> 50
accgctcagg tcattgacgt agttcatt 29

<210> 51
<211> 11
<212> DNA
<213> Artificial Sequence

<220>
<223> 0 starting fragment, position 1

<400> 51
ggggggggaa a 11

<210> 52
<211> 11
<212> DNA
<213> Artificial Sequence

<220>
<223> 0 starting fragment, position 2

<400> 52
ggggggggaa c 11

<210> 53
<211> 12
<212> DNA
<213> Artificial Sequence

<220>
<223> 0 starting fragment, position 2

<400> 53
ccccccccc tt 12

<210> 54
<211> 10
<212> DNA
<213> Artificial Sequence

<220>
<223> 1 starting fragment, position 2

<400> 54
aaaaaaaaac 10

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<210> 55
<211> 11
<212> DNA
<213> Artificial Sequence

<220>
<223> 0 starting fragment, position 7

<400> 55
ggggggggcc g 11

<210> 56
<211> 12
<212> DNA
<213> Artificial Sequence

<220>
<223> 0 starting fragment, position 7

<400> 56
cccccccccg cg 12

<210> 57
<211> 10
<212> DNA
<213> Artificial Sequence

<220>
<223> 1 starting fragment, position 7

<400> 57
aaaaaaaaacg 10

<210> 58
<211> 11
<212> DNA
<213> Artificial Sequence

<220>
<223> 1 starting fragment, position 7

<400> 58
ttttttttgc g 11

<210> 59
<211> 12
<212> DNA
<213> Artificial Sequence

<220>
<223> 0 starting fragment, position 8

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<400> 59
cccccccccc gg 12

<210> 60
<211> 11
<212> DNA
<213> Artificial Sequence

<220>
<223> 1 starting fragment, position 8

<400> 60
tttttttcg g 11

<210> 61
<211> 14
<212> DNA
<213> Artificial Sequence

<220>
<223> Fragment 0, position 1.2

<400> 61
aaaggggggg gaaa 14

<210> 62
<211> 13
<212> DNA
<213> Artificial Sequence

<220>
<223> Fragment 1, position 1.3

<400> 62
aacaaaaaaaaa aaa 13

<210> 63
<211> 14
<212> DNA
<213> Artificial Sequence

<220>
<223> Fragment 0, position 8.1

<400> 63
tttccccccc cccg 14

<210> 64
<211> 13

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<212> DNA
 <213> Artificial Sequence

 <220>
 <223> Fragment 1, position 8.1

 <400> 64
 ttttttttt tcg

13

<210> 65
 <211> 14
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Fragment 0, position 8.2

 <400> 65
 gttccccccc cccg

14

<210> 66
 <211> 13
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Fragment 1, position 8.2

 <400> 66
 gtttttttt tcg

13

<210> 67
 <211> 14
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Fragment 0, position 8.3

 <400> 67
 cttccccccc cccg

14

<210> 68
 <211> 13
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Fragment 1, position 8.3

 <400> 68

cttttttt tcg

13

<210> 69
 <211> 31
 <212> DNA
 <213> Artificial Sequence

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 <220>
 <221> misc_feature
 <222> (8)..(13)
 <223> N is any nucleotide.

<400> 69
 catccacnng agntggactc atggtcgctg c

31

<210> 70
 <211> 32
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Initiation linker

 <220>
 <221> misc_feature
 <222> (1)..(14)
 <223> N is any nucleotide.

<400> 70
 ncatccacnn gagntggact catggtcgct gc

32

<210> 71
 <211> 33
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Initiation linker

 <220>
 <221> misc_feature
 <222> (1)..(15)
 <223> N is any nucleotide.

<400> 71
 nncatccacn ngagntggac tcatggtcgc tgc

33

<210> 72
 <211> 34
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Initiation linker

<220>
 <221> misc_feature
 <222> (1)..(16)
 <223> N is any nucleotide.

<400> 72
 nnnatccac nngagntgga ctcatggtcg ctgc

34

<210> 73
 <211> 35
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Initiation linker

<220>
 <221> misc_feature
 <222> (12)..(17)
 <223> N is any nucleotide.

<400> 73
 gcgtcatcca cnngagntgg actcatggtc gctgc

35

<210> 74
 <211> 36
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 <213> Artificial Sequence

<220>
 <223> Initiation linker

<220>
 <221> misc_feature
 <222> (1)..(18)
 <223> N is any nucleotide.

<400> 74
 ngcgtcatcc acnngagntg gactcatggt cgctgc

36

<210> 75
 <211> 37
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Initiation linker

<220>
 <221> misc_feature
 <222> (1)..(19)
 <223> N is any nucleotide.

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 nngcgtcatc cacnngagt ggactcatgg tcgctgc

37

<210> 76
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 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Initiation linker

<220>
 <221> misc_feature
 <222> (1)..(20)
 <223> N is any nucleotide.

<400> 76
 nnngcgtcat ccacnngagn tggactcatg gtcgctgc

38

<210> 77
 <211> 39
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Initiation linker

<220>
 <221> misc_feature
 <222> (1)..(21)
 <223> N is any nucleotide.

<400> 77
 nnnngcgtca tccacnngag ntggactcat ggtcgctgc

39

<210> 78
 <211> 40
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Initiation linker

<220>
 <221> misc_feature
 <222> (1)..(22)
 <223> N is any nucleotide.

<400> 78
 nnnnngcgtc atccacnnga gntggactca tggtcgctgc

40

<210> 79
 <211> 10
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Propagation linker HgaI

<220>
 <221> misc_feature
 <222> (1)..(5)
 <223> N is any nucleotide.

<400> 79
 nnnnngcgtc

10

<210> 80
 <211> 32
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Gene A from PHIX174

<400> 80
 gctggaggcc tccactatga aatcgctag ag

32

<210> 81
 <211> 28
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Gene A from PHIX174

<400> 81
ctggcgaaa atgagaaaat tcgaccta

28

<210> 82
<211> 13
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<213> Artificial Sequence

<220>
<223> Recognition motif of the N-terminal part of the hsdS subunit of StyR 1241

<220>
<221> misc_feature
<222> (4)..(9)
<223> N is any nucleotide.

<400> 82
gaannnnnnr tcg

13

<210> 83
<211> 14
<212> DNA
<213> Artificial Sequence

<220>
<223> Recognition motif of the C-terminal part of the hsdS subunit of StyR 1241

<220>
<221> misc_feature
<222> (4)..(10)
<223> N is any nucleotide.

<400> 83
tcannnnnnn rttc

14

<210> 84
<211> 13
<212> DNA
<213> Artificial Sequence

<220>
<223> Recognition motif of a new enzyme made by merging the N- and C-terminal parts of the hsdS subunit of StyR 1241

<220>
<221> misc_feature
<222> (4)..(9)

<223> N is any nucleotide.

<400> 84
gaannnnnnr ttc

13

<210> 85
<211> 40
<212> DNA
<213> Artificial Sequence

<220>
<223> Ligated initiation linker

<220>
<221> misc_feature
<222> (1)..(22)
<223> N is any nucleotide with the proviso that the sequence from 1 to 6 is complementary to the sequence from 40 to 35 of SEQ ID NO: 15

<400> 85
nnnnnnnnnc atccacnnga gntggactca tggtcgctgc

40

<210> 86
<211> 47
<212> DNA
<213> Artificial Sequence

<220>
<223> An example of sequences that generate 5'-4 base overhangs by BbsI and Esp3I

<220>
<221> misc_feature
<222> (1)..(47)
<223> N is any nucleotide.

<400> 86
nnnnnnnnnga gcngagacgn nnnnngaaga cnngagcnnn nnnnnnn

47

<210> 87
<211> 47
<212> DNA
<213> Artificial Sequence

<220>
<223> An example of sequences that generate 5'-4 base overhangs by BbsI and Esp3I

<220>
 <221> misc_feature
 <222> (1)..(47)
 <223> N is any nucleotide.

<400> 87
 nnnnnnnnnn gctcnngtct tcnnnnnnncg tctcngctcn nnnnnnnn

47

<210> 88
 <211> 29
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> An example of 5' -4 base overhangs generated by BbsI and Esp3I cleavage

<220>
 <221> misc_feature
 <222> (5)..(25)
 <223> N is any nucleotide.

<400> 88
 gagcngagac gnnnnnnngaa gacnngagc

29

<210> 89
 <211> 25
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> An example of 5' -4 base overhangs generated by BbsI and Esp3I cleavage

<220>
 <221> misc_feature
 <222> (5)..(25)
 <223> N is any nucleotide.

<400> 89
 gctcnngtct tcnnnnnnncg tctcn

25

<210> 90
 <211> 22
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> An example of ligation products between 5' -4 base overhangs generated

rated by BbsI and Esp3I cleavage

<220>
<221> misc_feature
<222> (1)..(22)
<223> N is any nucleotide.

<400> 90
nnnnnnnnga gcnnnnnnnn nn

22

<210> 91
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> An example of ligation products between 5' -4 base overhangs gene
rated by BbsI and Esp3I cleavage

<220>
<221> misc_feature
<222> (1)..(22)
<223> N is any nucleotide.

<400> 91
nnnnnnnnnnn gctcnnnnnn nn

22

<210> 92
<211> 51
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<213> Artificial Sequence

<220>
<223> An example of sequences that generate two 3' 3 base overhangs by
BsaXI

<220>
<221> misc_feature
<222> (1)..(51)
<223> N is any nucleotide.

<400> 92
nnnnnnnnga gnnnnnnnnn acnnnnnctc cnnnnnnnga gnnnnnnnnn n

51

<210> 93
<211> 51
<212> DNA
<213> Artificial Sequence

<220>
 <223> An example of sequences that generate two 3' 3 base overhangs by BsaXI

<220>
 <221> misc_feature
 <222> (1)..(51)
 <223> N is any nucleotide.

<400> 93
 nnnnnnnnnn ctcnnnnnnn ggagnnnnng tnnnnnnnnn ctcnnnnnnn n 51

<210> 94
 <211> 30
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 <213> Artificial Sequence

<220>
 <223> An example of 3' 3 base overhangs generated by BsaXI cleavage

<220>
 <221> misc_feature
 <222> (1)..(27)
 <223> N is any nucleotide.

<400> 94
 nnnnnnnnna cnnnnnctcc nnnnnnnnag 30

<210> 95
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 <212> DNA
 <213> Artificial Sequence

<220>
 <223> An example of 3' 3 base overhangs generated by BsaXI cleavage

<220>
 <221> misc_feature
 <222> (1)..(27)
 <223> N is any nucleotide.

<400> 95
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<210> 96
 <211> 44
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> An example of sequences that generated blunt ends by MlyI
 <220>
 <221> misc_feature
 <222> (1)..(44)
 <223> N is any nucleotide.

<400> 96
 nnnnnnnnnn nnnnnnnnnn nnnngagtcn nnnnnnnnnn nnnn

44

<210> 97
 <211> 26
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> An example of 3' 3 base overhangs generated by MlyI cleavage

<220>
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 <222> (1)..(26)
 <223> N is any nucleotide.

<400> 97
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26

<210> 98
 <211> 30
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Gene A from PHIX174

<400> 98
 ctacgcgatt tcatagtgga ggcctccagc

30

<210> 99
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 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Gene A from PHIX174

<400> 99
 ggtcgaattt tctcattttc cgccagca

28

<210> 100
<211> 10
<212> DNA
<213> Artificial Sequence

<220>
<223> 1 starting fragment, position 1

<400> 100
aaaaaaaaaa 10

<210> 101
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<212> DNA
<213> Artificial Sequence

<220>
<223> 1 starting fragment, position 2

<400> 101
ttttttttt t 11

<210> 102
<211> 13
<212> DNA
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<220>
<223> Fragment 1, position 1.2

<400> 102
aaaaaaaaaa aaa 13

<210> 103
<211> 14
<212> DNA
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<220>
<223> Fragment 0, position 1.3

<400> 103
aacggggggg gaaa 14

<210> 104
<211> 14
<212> DNA
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<220>
<223> Fragment 0, position 8.3

<400> 104
cttccccccc cccg

14

<210> 105
<211> 13
<212> DNA
<213> Artificial Sequence

<220>
<223> Fragment 1, position 8.3

<400> 105
ctttttttt tcg

13